

Amendments to the Claims

This listing of claims will replace all prior versions and listings of claims.

1 - 27. Cancelled.

<sup>1</sup>  
~~28.~~ (Previously Presented) An isolated polypeptide comprising an amino acid sequence selected from the group consisting of:

- (a) amino acids 1 to 331 of SEQ ID NO:2;
- (b) the complete amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. 97343;
- (c) amino acids 2 to 331 of SEQ ID NO:2;
- (d) the complete amino acid sequence excepting the N-terminal methionine encoded by the cDNA clone contained in ATCC Deposit No. 97343;
- (e) amino acids 24 to 331 of SEQ ID NO:2;
- (f) amino acids 27 to 331 of SEQ ID NO:2; and
- (g) the amino acid sequence of the mature form of NAF-1 encoded by the cDNA clone contained in ATCC Deposit No. 97343.

<sup>1</sup>  
<sup>2</sup>~~29.~~ (Previously Presented) The isolated polypeptide of claim ~~28~~<sup>g</sup> wherein said amino acid sequence is (a).

<sup>1</sup>  
<sup>3</sup>~~30.~~ (Previously Presented) The isolated polypeptide of claim ~~28~~<sup>g</sup> wherein said amino acid sequence is (b).

<sup>1</sup>  
<sup>4</sup>~~31.~~ (Previously Presented) The isolated polypeptide of claim ~~28~~<sup>g</sup> wherein said amino acid sequence is (c).

<sup>1</sup>  
<sup>5</sup>~~32.~~ (Previously Presented) The isolated polypeptide of claim ~~28~~<sup>g</sup> wherein said amino acid sequence is (d).

<sup>6</sup> ~~33~~ (Previously Presented) The isolated polypeptide of claim <sup>1</sup> ~~28~~ wherein said amino acid sequence is (e).

<sup>7</sup> ~~34~~ (Previously Presented) The isolated polypeptide of claim <sup>1</sup> ~~28~~ wherein said amino acid sequence is (f).

<sup>8</sup> ~~35~~ (Previously Presented) The isolated polypeptide of claim <sup>1</sup> ~~28~~ wherein said amino acid sequence is (g).

36 - 67. (Cancelled).

<sup>9</sup> ~~68~~ (New) An epitope bearing fragment of the human NAF-1 polypeptide selected from the group consisting of:

(a) an epitope bearing fragment consisting of at least amino acids 75 to 100 of SEQ ID NO:2;

(b) an epitope bearing fragment consisting of at least amino acids 168 to 180 of SEQ ID NO:2;

(c) an epitope bearing fragment consisting of at least amino acids 204 to 226 of SEQ ID NO:2;

(d) an epitope bearing fragment consisting of at least amino acids 258 to 281 of SEQ ID NO:2; and,

(e) an epitope bearing fragment consisting of at least amino acids 291 to 327 of SEQ ID NO:2.

<sup>10</sup> ~~69~~ (New) The epitope bearing fragment of claim <sup>9</sup> ~~68~~ wherein said fragment is (a).

<sup>11</sup> ~~70~~ (New) The epitope bearing fragment of claim <sup>9</sup> ~~68~~ wherein said fragment is (b).

<sup>12</sup> ~~71~~ (New) The epitope bearing fragment of claim <sup>9</sup> ~~68~~ wherein said fragment is (c).

<sup>13</sup>  
~~72~~ (New) The epitope bearing fragment of claim <sup>9</sup>~~68~~ wherein said fragment is  
(d). ^

<sup>14</sup>  
~~73~~ (New) The epitope bearing fragment of claim <sup>9</sup>~~68~~ wherein said fragment is  
(e). ^